

3 ' 5 ' Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PUTET WEIGHT LOSS DISEASE (WD)

Inventors: Jestin et al. Appl. No.: 09/514,245

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TECH CENTER 1600/2900

Leu Ala Ser Arg Cys Arg Cys Cys Arg Pro Leu Thr Leu Ser Phe Ala Leu Cys Trp Arg Val Glu Ala Ala Ala Ala Gly Arg Cys Arg *** His Phe His Trp Ala Gly Ala Cys Lys Pro Leu Pro Leu Val Glu Ala Ala Asp Thr Phe Ile Gly Leu TGG TCG CGT GAA GCC GTC GCC GTC GTG GAG CCG TCG CAG TCA CTT TTA CGG TTC 9 18 27 36 45 54 ACC AGC GCA CTT CGG CAG CGG CAG CAC CTC GGC AGC GTC AGT GAA AAT GCC AAG Thr Ser Ala Leu Arg Gln Arg Gln His Leu Gly Ser Val Ser Glu Asn Ala Lys Pro Ala His Phe Gly Ser Gly Ser Thr Ser Ala Ala Ser Val Lys Met Pro Ser Gln Arg Thr Ser Ala Ala Ala Ala Ala Pro Arg Gln Arg Gln *** Lys Cys Gln Ala

Ser Phe Arg Gly Ala Val Gly Tyr Ser Thr Pro Thr *** Gly *** Tyr Asp Lys Leu Phe Ala Ala Arg Leu Gly Met Leu Pro Pro His Glu Gly Lys Ile Ile Arg Leu Phe Leu Pro Gly Cys Gly Trp Leu Leu His Thr Asn Val Arg Leu Leu Gly GTT CTT TTC GCC GGG CGT TGG GGT ATT CTC CAC CCA CAA GTG GGA ATT ATT AGG 63 72 81 90 99 108 CAA GAA AAG CGG CCC GCA ACC CCA TAA GAG GTG GGT GTT CAC CCT TAA TAA TCC Gln Glu Lys Arg Pro Ala Thr Pro *** Glu Val Gly Val His Pro *** *** Ser Lys Lys Ser Gly Pro Gln Pro His Lys Arg Trp Val Phe Thr Leu Asn Asn Pro Arg Lys Ala Ala Arg Asn Pro Ile Arg Gly Gly Cys Ser Pro Leu Ile Ile Leu

Arg Pro Pro Ser Phe Cys Phe Val Pro Ala Glu Leu Arg Gly Lys Gln Asn Asn Gly Leu Leu Leu Phe Val Phe Tyr Pro Leu Lys Trp Asp Gly Lys Lys Ile Ile Glu Ser Ser Ser Phe Phe Leu Ile Arg Ser Ser Gly Ile Glu Arg Lys Ser ***

AAG GCT CCT CCT CTT TTT GTT TTA TGC CCT CGA AGG TTA GAG GGA AAA ACT AAT 117 126 135 144 153 162

TTC CGA GGA GGA GAA AAA CAA AAT ACG GGA GCT TCC AAT CTC CCT TTT TGA TTA

Phe Arg Gly Gly Glu Lys Gln Asn Thr Gly Ala Ser Asn Leu Pro Phe *** Leu Ser Glu Glu Glu Lys Asn Lys Ile Arg Glu Leu Pro Ile Ser Leu Phe Asp Tyr Pro Arg Arg Arg Lys Thr Lys Tyr Gly Ser Phe Gln Ser Pro Phe Leu Ile Ile

Gln Lys His Arg Pro Leu Asn Pro Leu Pro Tyr Phe Glu Glu Gly Gly Pro Thr Lys Asn Thr Ala Leu Phe Thr Gln Phe Leu Thr Ser Ser Arg Val Glu Leu Pro Lys Thr Gln Pro Ser Ser Pro Lys Ser Ser Pro Leu Val Gly *** Arg Trp Pro AAA ACA AAC ACC GCT CCT TCC AAA CCT TCT CCC ATC TTG AGG AGT GGA GGT CCC 171 180 189 207 216
TTT TGT TTG TGG CGA GGA AGG TTT GGA AGA GGG TAG AAC TCC TCA CCT CCA GGG
Phe Cys Leu Trp Arg Gly Arg Phe Gly Arg Gly *** Asn Ser Ser Pro Pro Gly Phe Val Cys Gly Glu Glu Gly Leu Glu Gly Arg Thr Pro His Leu Gln Gly Leu Phe Val Ala Arg Lys Val Trp Lys Arg Val Glu Leu Leu Thr Ser Arg Gly

Gln Ser Asn Gln *** Ser Ala Ser Lys *** Cys Pro Ser Thr Thr Asn Gln His Lys Arg Ile Lys Ser Leu Leu Leu Ser Lys Val Leu His Leu Pro Ile Lys Thr Asn Ala Phe Lys Ala Leu Phe Cys Val Lys Leu Leu Thr Phe His Tyr Lys Pro CAA ACG CTT AAA ACG ATT CTT CGT CTG AAA ATT GTT CCA CTT CAC CAT AAA ACC 225 234 243 252 261 270 GTT TGC GAA TTT TGC TAA GAA GCA GAC TTT TAA CAA GGT GAA GTG CTA TTT TGG Val Cys Glu Phe Cys *** Glu Ala Asp Phe *** Gln Gly Glu Val Val Phe Trp Phe Ala Asn Phe Ala Lys Lys Gln Thr Phe Asn Lys Val Lys Trp Tyr Phe Gly Leu Arg Ile Leu Leu Arg Ser Arg Leu Leu Thr Arg *** Ser Gly Ile Leu Val



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Gly Ser Gly Cys Arg Ser Leu Ser Leu Phe Arg Gly Ala Ser Tyr Leu Ile Ser Gly Ala Ala Val Asp Leu Phe Arg Phe Ser Gly Val Leu Leu Ile Phe Phe Val Ala Arg Gln Trp Met Ser Phe Ala Phe Pro Val Ser Trp Cys Phe Leu Ser Tyr ACG GGC GAC GGT GTA GCT CTT TCG CTT TCC TTG GCT GGT CGT CTT ATT TCT TAT 279 288 297 306 315 324 TGC CCG CTG CCA CAT CGA GAA AGC GAA AGG AAC CGA CCA GCA GAA TAA AGA ATA Cys Pro Leu Pro His Arg Glu Ser Glu Arg Asn Arg Pro Ala Glu *** Arg Ile Ala Arg Cys His Ile Glu Lys Ala Lys Gly Thr Asp Gln Gln Asn Lys Glu Tyr Pro Ala Ala Thr Ser Arg Lys Arg Lys Glu Pro Thr Ser Arg Ile Lys Asn Thr

Cys Tyr Leu Leu Gly Cys Val *** Arg Thr His Leu Glu Ala Ser Gly Pro Ser Ala Thr Phe Phe Ala Val Tyr Lys Asp Leu Thr Ser Ser Arg Pro Val Leu Pro Gln Leu Leu Ser Pro Trp Met Ser Ile Ser His Pro Ala Gly Arg Phe Trp Pro GAC GTC ATT TCT TCC GGT GTA TGA ATA GCT CAC ACC TCG AGG CGC CTT GGT CCC 333 342 351 360 369 378 CTG CAG TAA AGA AGG CCA CAT ACT TAT CGA GTG TGG AGC TCC GCG GAA CCA GGG Leu Gln *** Arg Arg Pro His Thr Tyr Arg Val Trp Ser Ser Ala Glu Pro Gly Cys Ser Lys Glu Gly His Ile Leu Ile Glu Cys Gly Ala Pro Arg Asn Gln Gly Ala Val Lys Lys Ala Thr Tyr Leu Ser Ser Val Glu Leu Arg Gly Thr Arg Gly

Ala Cys Arg Gly Thr *** Gln Gln Ser Tyr Gly Lys Pro Ser Pro Thr Lys Pro Leu Ala Ala Val Gln Arg Ser Ser His Thr Gly Lys Gln Leu Arg Pro Arg Gln Phe Arg Leu Ser Arg Asp Val Ala Thr Leu Val Arg Lys Ser Val Pro Asp Lys CTT CGC GTC GCT GGA CAG ATG ACG ACA CTC ATG GGA AAA CCT CTG CCC CAG AAA 387 405 414 423 432 432 GAA GCG CAG CGA CCT GTC TAC TGC TGT GAG TAC CCT TTT GGA GAC GGG GTC TTT Glu Ala Gln Arg Pro Val Tyr Cys Cys Glu Tyr Pro Phe Gly Asp Gly Val Phe Lys Arg Ser Asp Leu Ser Thr Ala Val Ser Thr Leu Leu Glu Thr Gly Ser Leu Ser Ala Ala Thr Cys Leu Leu *** Val Pro Phe Trp Arg Arg Gly Leu Trp

Ser Gln Leu Arg Ala Thr Glu Gln Leu Thr His Ser Phe Asn Gly Arg Ala Pro His Ser Tyr Gly Leu Leu Lys Arg Tyr Arg Ile His Ser Ile Glu Ala Pro Gln Thr Val Thr Ala Ser Cys Asn Gly Thr Val Tyr Thr Leu Phe Lys Arg Pro Ser CCA CTG ACA TCG GCT CGT CAA AGG ACA TTG CAT ACA CTC TTT AAA GGC GCC CGA 441 450 459 468 477 486 GGT GAC TGT AGC CGA GCA GTT TCC TGT AAC GTA TGT GAG AAA TTT CCG CGG GCT Gly Asp Cys Ser Arg Ala Val Ser Cys Asn Val Cys Glu Lys Phe Pro Arg Ala Val Thr Val Ala Glu Gln Phe Pro Val Thr Tyr Val Arg Asn Phe Arg Gly Leu *** Leu *** Pro Ser Ser Phe Leu *** Arg Met *** Glu Ile Ser Ala Gly Trp

Gln Val Lys Ser Leu Ser Arg Ser Ser Ala Ala Ala His Asn Ser Ser Leu Gln Ser Phe Lys Gln Phe His Ala Pro Leu His Leu Leu Thr Ile Pro Leu Cys Ser Ala Ser Ser Lys Phe Thr Leu Pro Phe Ile Cys Cys Arg Ser Gln Phe Val Ala CCG ACT TGA AAA CTT TCA CTC GCC CTT CTA CGT CGT CGC ACT AAC CTT CTG TCG 495 513 540 GGC TGA ACT TTT GAA AGT GAG CGG GAA GAT GCA GCG TGA TTG GAA GAC AGC Gly *** Thr Phe Glu Ser Glu Arg Glu Asp Ala Ala Ala *** Leu Glu Asp Ser Ala Glu Leu Leu Lys Val Ser Gly Lys Met Gln Gln Arg Asp Trp Lys Thr Ala Leu Asn Phe *** Lys *** Ala Gly Arg Cys Ser Ser Val Ile Gly Arg Gln Leu



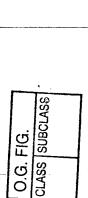
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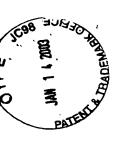
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DRAFTSN

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Val Arg *** Leu Pro Gly Ala Arg Asn His Ser Ser Gly Thr Pro Gly Tyr Asn Tyr Val Asp Tyr His Ala Arg Gly Thr Thr Pro Leu Ala Leu Pro Gly Thr Ile Thr Cys Thr Met Thr Pro Gly Gly Pro Gln Pro Phe Leu Trp His Ala Arg Leu ACA TGT GCA GTA TCA CCC GGG CGG GCC AAC ACC CTT CTC GGT CAC CCG GGC ATT 549 558 567 576 585 594 TGT ACA CGT CAT AGT GGG CCC GCC CGG TTG TGG GAA GAG CCA GTG GGC CCG TAA Cys Thr Arg His Ser Gly Pro Ala Arg Leu Trp Glu Glu Pro Val Gly Pro ***
Val His Val Ile Val Gly Pro Pro Gly Cys Gly Lys Ser Gln Trp Ala Arg Asn
Tyr Thr Ser *** Trp Ala Arg Pro Val Val Gly Arg Ala Ser Gly Pro Val Ile Gln Gln Ala *** Pro Cys Arg Ser Ser Ala *** Tyr Phe Tyr Thr Thr Pro His Lys Ser Leu Arg Pro Val Gly Val Pro Leu Arg Thr Ser Ile Leu Pro Pro Ile Lys Ala Ser Gly Leu Ser Val *** Gln Phe Gly Leu Leu Phe Leu His His Ser AAA ACG ACT CGG ATC CCT GTG GAT GAC CTT CGG ATC ATC TTT ATT CAC CAC CCT 603 621 630 639 TTT TGC TGA GCC TAG GGA CAC CTA CTG GAA GCC TAG TAG AAA TAA GTG GTG GGA Phe Cys *** Ala *** Gly His Leu Leu Glu Ala *** *** Lys *** Val Val Gly Phe Ala Glu Pro Arg Asp Thr Tyr Trp Lys Pro Ser Arg Asn Lys Trp Trp Asp Leu Leu Ser Leu Gly Thr Pro Thr Gly Ser Leu Val Glu Ile Ser Gly Gly Met Ile Asp His Leu Leu Leu Gln Gln Lys Pro His Asn Lys His Ser Thr Val Lys Ser Ile Met Ser Phe Phe Asn Asn Gln Ile Ile Lys Ile Ala Pro *** Arg Pro Tyr *** Pro Ser Ser Thr Thr Thr Lys Ser Ser Lys *** Pro Gln Asn Gly ACC TAT AGT ACC TCT TCT TCA ACA ACA AAA CCT ACT AAA AAT ACC GAC CAA TGG 657 TGG ATA TCA TGG AGA AGA AGT TGT TGT TTT GGA TGA TTT TTA TGG CTG GTT ACC Trp Ile Ser Trp Arg Arg Ser Cys Cys Phe Gly *** Phe Leu Trp Leu Val Thr Gly Tyr His Gly Glu Glu Val Val Leu Asp Asp Phe Tyr Gly Trp Leu Pro Asp Ile Met Glu Lys Lys Leu Leu Phe Trp Met Ile Phe Met Ala Gly Tyr Leu Pro His Asp Val Ser Val Thr His Gly Thr Asp Met Ser Gln Leu Ser *** Leu Pro Ile Ile *** Gln Ser Gln Thr Val Pro Ile Trp Gln Ser Tyr Leu Ser Phe Gln Ser Ser Arg Ser Leu Ser His Ser Arg Tyr Gly Asn Val Thr Ser Val Leu AAC CCT ACT AGA TGA CTC TGA CAC ACT GGC CAT AGG TAA CTG ACA TCT CTG ATT 756 729 TTG GGA TGA TCT ACT GAG ACT GTG TGA CCG GTA TCC ATT GAC TGT AGA GAC TAA Leu Gly *** Ser Thr Glu Thr Val *** Pro Val Ser Ile Asp Cys Arg Asp *** Trp Asp Asp Leu Leu Arg Leu Cys Asp Arg Tyr Pro Leu Thr Val Giu Thr Lys Gly Met Ile Tyr *** Asp Cys Val Thr Gly Ile His *** Leu *** Arg Leu Lys Pro Tyr Gln Glu Lys Lys Pro Gly Cys Tyr Lys Ser *** Trp Cys Asp Pro Gly Pro Thr Ser Asn Arg Lys Gln Gly Ala Thr Asn Gln Asn Gly Ala Ile Leu Gly Pro Pro Val Thr Gly Lys Lys Ala Arg Leu Ile Lys Ile Val Leu Leu *** Ala TCC CCC ATG ACA AGG AAA AAA CCG GGC GTC ATA AAA CTA ATG GTC GTT AGT CCG AGG GGG TAC TGT TCC TTT TTT ĞGC CCG CAG TAT TTT GAT TAC CAG CAA TCA GGC Arg Gly Tyr Cys Ser Phe Phe Gly Pro Gln Tyr Phe Asp Tyr Gln Gln Ser Gly Gly Thr Val Pro Phe Leu Ala Arg Ser Ile Leu Ile Thr Ser Asn Gln Ala Gly Val Leu Phe Leu Phe Trp-Pro-Ala Val-Phe *** Leu Pro Ala Ile Arg Pro



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Gly Pro Ile Thr Ser Arg Leu Gln Gln Gly Leu Gln Leu Leu Glu Arg Asp Ser Gly Leu Phe Pro Val Gly *** Ser Ser Asp Trp Ser Tyr Phe Ser Glu Ile Pro Gly Trp Ser His Tyr Glu Glu Val Ala Thr Gly Ala Thr Ser Ala Arg *** Arg GGG GGT CCT TAC CAT GAG GAG TTG ACG ACA GGG TCG ACA TCT TCG AGA GAT AGC 819 828 837 846 855 864 CCC CCA GGA ATG GTA CTC CTC AAC TGC TGT CCC AGC TGT AGA AGC TCT CTA TCG Pro Pro Gly Met Val Leu Leu Asn Cys Cys Pro Ser Cys Arg Ser Ser Leu Ser Pro Gln Glu Trp Tyr Ser Ser Thr Ala Val Pro Ala Val Glu Ala Leu Tyr Arg Pro Arg Asn Gly Thr Pro Gln Leu Leu Ser Gln Leu *** Lys Leu Ser Ile Gly

Ser *** *** Lys Ala Ile Lys Ser Ser Gln Gln Leu Val Ile Trp Pro Pro Val Pro Asn Ser Ser Gln Leu Lys Pro Leu Ser Ser Ser Phe Leu Gly Arg Leu Tyr Leu Ile Val Val Lys Cys Asn Gln Phe Val Ala Pro Ser Cys Asp Val Ser Thr CTC CTA ATG ATG AAA CGT TAA AAC CTT CTG ACG ACC TCT TGT TAG GTG CCT CCA 873 882 891 900 918 GAG GAT TAC TAC TTT GCA ATT TTG GAA GAC TGC TGG AGA ACA ATC CAC GGA GGT Glu Asp Tyr Tyr Phe Ala Ile Leu Glu Asp Cys Trp Arg Thr Ile His Gly Gly Arg Ile Thr Thr Leu Gln Phe Trp Lys Thr Ala Gly Glu Gln Ser Thr Glu Val Gly Leu Leu Leu Cys Asn Phe Gly Arg Leu Leu Glu Asn Asn Pro Arg Tyr

Arg Leu Gly Ile Gln Leu Leu Pro Gly Val Arg His Gly Lys Gly Met Tyr Phe Gly Phe Ala Ser Lys Phe Cys His Val Trp Gly Thr Gly Lys Glu Trp Ile Phe Gly Ser Pro Arg Asn Ser Ala Thr Ser Gly Gly Gln Ala Arg Lys Gly Tyr Leu

TGG GCT TCC GGC TAA ACT TCG TCA CCT GGG TGG GAC ACG GGA AAA GGG TAT ATT 927 936 945 954 963 972

ACC CGA AGG CCG ATT TGA AGC AGT GGA CCC ACC CTG TGC CCT TTT CCC ATA TAA

Thr Arg Arg Pro Ile *** Ser Ser Gly Pro Thr Leu Cys Pro Phe Pro Ile ***

Pro Glu Gly Arg Phe Glu Ala Val Asp Pro Pro Cys Ala Leu Phe Pro Tyr Lys Pro Lys Ala Asp Leu Lys Gln Trp Thr His Pro Val Pro Phe Ser His Ile Lys

Leu Asn Ser Leu Arg Lys Gln *** *** Met Thr Ile Thr Lys Ile Lys Ile ***
Tyr Ile Val Ser Asp Lys Lys Asn Asp Cys Arg Leu Pro Lys *** Lys *** Glu
Ile Phe *** Gln Thr Lys Lys Thr Ile Val Asp Tyr His Asn Lys Asn Lys Asn
TTA TTT AAT GAC TCA GAA AAA ACA ATA GTG TAG CAT TAC CAA AAA TAA AAA TAA
981
990
999
1008
1017
1026
AAT AAA TTA CTG AGT CTT TTT TGT TAT CAC ATC GTA ATG GTT TTT ATT TTT ATT
Asn Lys Leu Leu Ser Leu Phe Cys Tyr His Ile Val Met Val Phe Ile Phe Ile
Ile Asn Tyr *** Val Phe Phe Val Ile Thr Ser *** Trp Phe Leu Phe Leu Phe
*** Ile Thr Glu Ser Phe Leu Leu Ser His Arg Asn Gly Phe Tyr Phe Tyr Ser

Lys Ser Pro Arg Glu Pro Tyr Ile Arg Gln Ile Thr Cys Leu Tyr Asp Val Lys Asn Leu Pro Asp Lys Leu Ile Phe Glu Arg Phe Gln Val Tyr Ile Thr Leu Arg Met *** Leu Thr Lys *** Ser Leu Asn Glu Ser Asn Tyr Met Phe Leu *** Gly GTA AAT CTC CCA GAA AGT CCT ATT TAA GAG ACT TAA CAT GTA TTT ATC AGT TGG 1035 1044 1053 1062 1071 1080 CAT TTA GAG GGT CTT TCA GGA TAA ATT CTC TGA ATT GTA CAT AAA TAG TCA ACC His Leu Glu Gly Leu Ser Gly *** Ile Leu *** Ile Val His Lys *** Ser Thr Ile *** Arg Val Phe Gln Asp Lys Phe Ser Glu Leu Tyr Ile Asn Ser Gln Pro Phe Arg Gly Ser Phe Arg Ile Asn Ser Leu Asn Cys Thr *** Ile Val Asn Leu

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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWI

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SUBCLASS EG. CLASS ග් Ö 应

Gly Cys Leu Lys Pro Ser His Asn Cys Lys Pro Ala Cys Leu Gly Pro Arg His Val Val Tyr Asn Gln Ala Thr Thr Ala Asn Gln Leu Ala Tyr Gly Leu Gly Thr *** Trp Met Ile Lys Pro Gln Pro Gln Met Lys Ser Arg Met Ala Trp Ala Gln AAT GGT_GTA_TTA_AAA_CCC GAC_ACC_AAC_GTA_AAA_CCT_CGC_GTA_TCG_GGT_CCG_GAC_1089 1098 1107 1116 1125 1134 TTA CCA CAT AAT TTT GGG CTG TGG TTG CAT TTT GGA GCG CAT AGC CCA GGC CTG Leu Pro His Asn Phe Gly Leu Trp Leu His Phe Gly Ala His Ser Pro Gly Leu Tyr His Ile Ile Leu Gly Cys Gly Cys Ile Leu Glu Arg Ile Ala Gln Ala Cys Thr Thr *** Phe Trp Ala Val Val Ala Phe Trp Ser Ala *** Pro Arg Pro Val

Ala Arg Cys Gln His Pro Tyr Lys Phe Pro Ala Val Ala Pro Lys Lys *** ***
His Glu Val Asn Thr His Thr Asn Leu His Leu Trp Leu Gln Asn Arg Lys Asn
Thr Ser Ser Met Pro Thr Pro Ile *** Ile Ser Gly Cys Ser Thr Glu Lys Ile ACA CGA GCT GTA ACC ACA CCC ATA AAT TTA CCT CGG TGT CGA CCA AAG AAA ATA 1143 1152 1161 1170 1179 1188 TGT GCT CGA CAT TGG TGT GGG TAT TTA AAT GGA GCC ACA GCT GGT TTC TTT TAT Cys Ala Arg His Trp Cys Gly Tyr Leu Asn Gly Ala Thr Ala Gly Phe Phe Tyr Val Leu Asp Ile Gly Val Gly Ile *** Met Glu Pro Gln Leu Val Ser Phe Ile Cys Ser Thr Leu Val Trp Val Phe Lys Trp Ser His Ser Trp Phe Leu Leu

Lys Ala Pro Val Leu *** Asn Asn Pro Arg Ala Arg Thr Gln Pro His Leu Val Asn Pro Gln Phe Trp Asp Ile Thr Gln Asp Leu Glu Pro Lys Pro Thr Phe Tyr Ile Gln Ser Ser Gly Ile Leu Gln Lys Thr *** Ser Gln Asn Pro Pro Ser Thr ATA AAC CGA CCT TGG TTA GTT AAC AAA CCA GAT CGA GAC CAA ACC CCC ACT TCA 1197 1206 1215 1224 1233 1242 TAT TTG GCT GGA ACC AAT CAA TTG TTT GGT CTA GCT CTG GTT TGG GGG TGA AGT Tyr Leu Ala Gly Thr Asn Gln Leu Phe Gly Leu Ala Leu Val Trp Gly *** Ser Ile Trp Leu Glu Pro Ile Asn Cys Leu Val *** Leu Trp Phe Gly Gly Gly Val Phe Gly Trp Asn Gln Ser Ile Val Trp Ser Ser Gly Leu Gly Val Lys Tyr

Gln Leu Pro Leu Tyr Leu Ala Ala Lys His His Pro Pro Leu Leu *** Tyr Arg Ser His Tyr Thr Phe Pro Gln Arg Ile Thr His Arg Ser Ser Tyr Asn Ile Gly Pro Thr Thr Pro Leu Pro Ser Gly *** Pro Thr Ala Pro Pro Thr Thr Leu TGG ACC TCA CCA TCC ATT TCC CGA CGG AAT ACC ACA CCG CCC TCC TCA TCA ATT 1251 1260 1269 1278 1287 1296 ACC TGG AGT GGT AGG TAA AGG GCT GCC TTA TGG TGT GGC GGG AGG AGT AGT TAA Thr Trp Ser Gly Arg *** Arg Ala Ala Leu Trp Cys Gly Gly Arg Ser Ser ***
Pro Gly Val Val Gly Lys Gly Leu Pro Tyr Gly Val Ala Gly Gly Val Val Asn
Leu Glu Trp *** Val Lys Gly Cys Leu Met Val Trp Arg Glu Glu *** Leu Ile

Leu Pro *** Leu Gly Leu Gln His Leu Pro Asn Cys Leu Gln Cys Gly Leu Tyr Tyr Pro Asp Tyr Ala Leu Asn Thr Ser Pro Thr Val Phe Asn Ala Asp Leu Ile Ile Pro Thr Met Pro Trp Thr Pro Pro Pro Pro *** Leu Thr Pro Met Trp Ser ATA TCC CCA GTA TCC GGT TCA ACC ACC TCC CCC AAT GTT TCA ACC GTA GGT TCT 1305 1314 1323 1332 1341 1350 TAT AGG GGT CAT AGG CCA AGT TGG TGG AGG GGG TTA CAA AGT TGG CAT CCA AGA Tyr Arg Gly His Arg Pro Ser Trp Trp Arg Gly Leu Gln Ser Trp His Pro Arg Ile Gly Val Ile Gly Gln Val Gly Gly Gly Tyr Lys Val Gly Ile Gln Asp *** Gly Ser *** Ala Lys Leu Val Glu Gly Val Thr Lys Leu Ala Ser Lys Ile



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Cys Cys His Val Trp Cys Arg Lys Ser *** Leu His His Pro Arg Gln Pro Leu Val Val Thr Ser Gly Val Gly Arg Gln Asn Ser Thr Ile Pro Asp Arg Pro Tyr Leu Leu Pro Gly Leu Val Glu Lys Ile Leu Pro Ser Pro Thr Glu Pro Thr TAA CAA CĂĞ TGG ACC CĂĂ CAC CTC TTT GAT TAG ĂĞĞ TGA TGG ĞĞT CTC TGG ĞĞT Gln Gln Trp Thr Gln His Leu Phe Asp *** Arg *** Trp Gly Leu Trp Gly Asn Asn Ser Gly Pro Asn Thr Ser Leu Ile Arg Gly Asp Gly Val Ser Gly Val Thr Thr Val Asp Pro Thr Pro Leu *** Leu Glu Val Met Gly Ser Leu Gly ***

Ile *** Ile *** Gly Lys *** Tyr Pro Leu Ile Pro Phe Thr Pro Thr Pro Pro Phe Glu Tyr Lys Ala Lys Arg Ile Arg Tyr Tyr Gln Phe Pro Leu Pro Phe Asn Met Asn Leu Arg Glu Leu Val Thr Thr Asn Ser Leu Tyr Pro Tyr Pro TTT TAA GTA TAA ATC GGA AAG ATT ATG CCA TCA TAA CCT TTC CAT CCC CAT CCC 1422 1440 1449 1458 1431 AAA ATT CAT ATT TAG CCT TTC TAA TAC GGT AGT ATT GGA AAG GTA GGG GTA GGG Lys Ile His Ile *** Pro Phe *** Tyr Gly Ser Ile Gly Lys Val Gly Val Gly
Lys Phe Ile Phe Ser Leu Ser Asn Thr Val Val Leu Glu Arg *** Gly *** Gly Asn Ser Tyr Leu Ala Phe Leu Ile Arg *** Tyr Trp Lys Gly Arg Gly Arg Gly

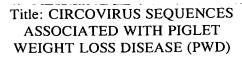
Gln His Arg Arg Leu Pro Pro Pro Val Pro Arg His Gln Ile Glu Ala Arg ***
Asn Thr Gly Gly Ser Pro Pro Leu Phe Gln Gly Ile Asn Phe Arg Leu Glu Asn
The Dro Ala Ala Clar Pro Pro Cor Cor Ala Cor The Cor Arg Leu Glu Asn Thr Pro Ala'Ala'Gln Pro Pro Ser Ser Ser Ala'Ser Thr Ser Asp ** Ser Thr CCA ACC ACG GCG GAC TGG CCC CCT CCT TGA CCG GCT ACA ACT TAG AGT CGA GCA 1467 1476 1485 1494 1503 1512 GGT TGG TGC CGC CTG AGG GGG GGA GGA ACT GGC CGA TGT TGA ATC TCA GCT CGT Gly Trp Cys Arg Leu Arg Gly Gly Gly Thr Gly Arg Cys *** Ile Ser Ala Arg Val Gly Ala Ala *** Gly Gly Glu Glu Leu Ala Asp Val Glu Ser Gln Leu Val Leu Val Pro Pro Glu Gly Gly Arg Asn Trp Pro Met Leu Asn Leu Ser Ser Leu

Cys Glu Leu Ile Ala Ala Leu Thr Arg Arg Lys His His Thr Cys Ile Arg *** Val Asn Trp Ser Pro Gln Ser His Gly Gly Arg Ile Thr Leu Val Phe Glu Arg Leu Met Gly Leu His Ser Arg Thr Asp Glu Glu *** Pro Ser Tyr Leu Asn Glu ATT GTA AGG TTC TAC CGA CGC TCA CAG GAG GAG AAT ACC ACT CAT GTT TAA GAG 1521 1530 1539 1548 1557 1566 TAA CAT TCC AAG ATG GCT GCG AGT GTC CTC CTC TTA TGG TGA GTA CAA ATT CTC *** His Ser Lys Met Ala Ala Ser Val Leu Leu Leu Trp *** Val Gln Ile Leu Asn Ile Pro Arg Trp Leu Arg Val Ser Ser Ser Tyr Gly Glu Tyr Lys Phe Ser Thr Phe Gln Asp Gly Cys Glu Cys Pro Pro Leu Met Val Ser Thr Asn Ser Leu

Phe Pro Pro Phe Gln Leu Tyr Gly Asp Lys Pro Ala Met Gln Leu Pro Lys Gln Ser Leu Arg Ser Asn Phe Ile Gly Thr Lys Arg Arg Trp Arg Tyr Arg Asn Arg Leu Phe Ala Pro Ile Ser Ser Val Arg Arg Glu Ala Gly Asp Thr Val Thr Glu ATC TTT CCG CCC TTA ACT TCT ATG GGC AGA AAG CCG CGG TAG ACA TTG CCA AAG 1575 1584 1593 1602 1611 1620 TAG AAA GGC GGG AAT TGA AGA TAC CCG TCT TTC GGC GCC ATC TGT AAC GGT TTC *** Lys Gly Gly Asn *** Arg Tyr Pro Ser Phe Gly Ala Ile Cys Asn Gly Phe Arg Lys Ala Gly Ile Glu Asp Thr Arg Leu Ser Ala Pro Ser Val Thr Val Ser Glu Arg Arg Glu Leu Lys Ile Pro Val Phe Arg Arg His Leu *** Arg Phe Leu

FIG. 2f





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APPROVED O.G. FIG.

BY CLASS SUBCLASS

RAFTSMAN

Leu Arg Pro Thr Gly Phe Ile Thr Lys Glu Pro Pro His Lys Trp Ser Pro Gln Phe Ala Pro His Val Leu Tyr Pro Arg Arg Arg Leu Ile Asn Gly Leu His Ser Ser Pro Pro Thr Tyr Trp Ile His Asp Glu Gly Ser Ser Thr Glu Leu Ile Ala

ACT TCC GCC CCA CAT GGT TTA TAC CAG AAG AGG CCT CCT ACA AAG GTT CTA CCG

1629
1638
1647
1656
1665
1674
TGA AGG CGG GGT GTA CCA AAT ATG GTC TTC TCC GGA GGA TGT TTC CAA GAT GGC

*** Arg Arg Gly Val Pro Asn Met Val Phe Ser Gly Gly Cys Phe Gln Asp Gly Glu Gly Gly Val Tyr Gln Ile Trp Ser Ser Pro Glu Asp Val Ser Lys Met Ala Lys Ala Gly Cys Thr Lys Tyr Gly Leu Leu Arg Arg Met Phe Pro Arg Trp Leu

Pro Pro Pro Asp Thr Lys Gln Pro Leu Ala Glu Lys Ala Val Asp Asp *** Leu Arg Pro Arg Thr Arg Arg Arg Arg Tyr Arg Arg Arg Pro Trp Thr Met Arg Tyr Ala Pro Ala Pro Gly Asp Glu Ala Thr Val Gly Gly Gln Gly Arg *** Gly Ile

ACG CCC CCG CCC AGG CAG AAG ACG CCA TTG CGG AGG AAC CGG TGC AGT AGG ATA

1683 1692 1701 1710 1719 1728

TGC GGG GGC GGG TCC GTC TTC TGC GGT AAC GCC TCC TTG GCC ACG TCA TCC TAT

Cys Gly Gly Gly Ser Val Phe Cys Gly Asn Ala Ser Leu Ala Thr Ser Ser Tyr Ala Gly Ala Gly Pro Ser Ser Ala Val Thr Pro Pro Trp Pro Arg His Pro Ile

Arg Gly Arg Val Arg Leu Leu Arg *** Arg Leu Leu Gly His Val Ile Leu ***

Leu Ser Leu Leu Ala Ser Ser Tyr Tyr
Phe His Phe Phe His Ala Ala Thr Thr Asn
Phe Thr Phe Ser Thr Arg Gln Gln Leu Ile

TTT TCA CTT TCT TCA CGC GAC GAC ATC ATA A 5'
1737
1746
AAA AGT GAA AGA AGT GCG CTG CTG TAG TAT T 3'

Lys Ser Glu Arg Ser Ala Leu Leu *** Tyr
Lys Val Lys Glu Val Arg Cys Cys Ser Ile
Lys *** Lys Lys Cys Cys Ala Ala Val Val

FIG. 2g



CLASS SUBCLASS

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circopormank circopormeeh circopordfp

APPROVED O.G. FIG.

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circopormank circopormeeh circopordfp	10 20 30 40 50 1 ACCAGCGCAC TTCGGCAGCG GCAGCACCTC GGCAGCGTCA GTGAAAATGC 1 ACCAGCGCAC TTCGGCAGCG GCAGCACCTC GGCAGCGTCA GTGAAAATGC 1 ACCAGCGCAC TTCGGCAGCG GCAGCACCTC GGCAGCGTCA GTGAAAATGC	50 50 50
circopormank circopormeeh circopordfp	60 70 80 90 100 51 CAAGCAAGAA AAGCGGCCCG CAACCCCATA AGAGGTGGGT GTTCACCCTT	100 100 100
circopormank circopormeeh circopordfp	110 120 130 140 150 101 [AATAATCCTT] [CCGAGGAGGA] [GAAAAACAAA] [ATACGGGAGC] [TTCCAATCTC] 101 [AATAATCCTT] [CCGAGGAGGA] [GAAAAACAAA] [ATACGGGAGC] [TTCCAATCTC] 101 [AATAATCCTT] [CCGAGGAGGA] [GAAAAACAAA] [ATACGGGAGC] [TTCCAATCTC]	150 150 150
circopormank circopormeeh circopordfp	160 170 180 190 200 151 CCTTTTTGAT TATTTTGTTT GCGGAGAGGA AGGTTTGGAA GAGGGTAGAA 151 CCTTTTTGAT TATTTTGTTT GCGGAGAGGA AGGTTTGGAA GAGGGTAGAA 151 CCTTTTTGAT TATTTTGTTT GTGGCGAGGA AGGTTTGGAA GAGGGTAGAA	200 200 200
circopormank circopormeeh circopordfp	210 220 230 240 250 201 CTCCTCACCT CCAGGGGTTT GCTAATTTTG CTAAGAAGCA GACTTTTAAC 201 CTCCTCACCT CCAGGGGTTT GCGAATTTTG CTAAGAAGCA GACTTTTAAC 201 CTCCTCACCT CCAGGGGTTT GCGAATTTTG CTAAGAAGCA GACTTTTAAC	250 250 250
circopormank circopormeeh circopordfp	260 270 280 290 300 251 AAGGTGAAGT GGTATTTTGG TGCCCGCTGC CACATCGAGA AAGCGAAAGG 251 AAGGTGAAGT GGTATTTTGG TGCCCGCTGC CACATCGAGA AAGCGAAAGG 251 AAGGTGAAGT GGTATTTTGG TGCCCGCTGC CACATCGAGA AAGCGAAAGG	300 300 300
circopormank circopormeeh circopordfp	310 320 330 340 350 301 [AACCGACCAG] [CAGAATAAAG] [AATACTGCAG] [TAAAGAAGGC] [CACATACTTA] 301 [AACCGACCAG] [CAGAATAAAG] [AATACTGCAG] [TAAAGAAGGC] [CACATACTTA] 301 [AACCGACCAG] [CAGAATAAAG] [AATACTGCAG] [TAAAGAAGGC] [CACATACTTA]	350 350 350
circopormank circopormeeh circopordfp	360 370 380 390 400 351 [TCGAGTGTGG] [AGCTCCGCGG] [AACCAGGGGA] [AGCGCAGCGA] [CCTGTCTACT] 351 [TCGAGTGTGG] [AGCTCCGCGG] [AACCAGGGGA] [AGCGCAGCGA] [CCTGTCTACT] 351 [TCGAGTGTGG] [AGCTCCGCGG] [AACCAGGGGA] [AGCGCAGCGA] [CCTGTCTACT]	400 400 400
circopormank circopormeeh circopordfp	410 420 430 440 450 401 [GCTGTGAGTA] [CCCTTTTGGA] [GACGGGGTCT] [TTGGTGACTG] [TAGCCGAGCA] 401 [GCTGTGAGTA] [CCCTTTTGGA] [GACGGGGTCT] [TTGGTGACTG] [TAGCCGAGCA] 401 [GCTGTGAGTA] [CCCTTTTGGA] [GACGGGGTCT] [TTGGTGACTG] [TAGCCGAGCA]	450 450 450
circopormank circopormeeh circopordfp	460 470 480 490 500 451 [GTTCCCTGTA] [ACGTATGTGA] [GAAATTTCCG] [CGGGCTGGCT] [GAACTTTTGA] 451 [GTTCCCTGTA] [ACGTATGTGA] [GAAATTTCCG] [CGGCTGGCT] [GAACTTTTGA] 451 [GTTCCTGTA] [ACGTATGTGA] [GAAATTTCCG] [CGGCTGGCT] [GAACTTTTGA]	500 500 500
circopormank circopormeeh circopordfp	510 520 530 540 550 501 AAGTGAGCGG GAAGATGCAG CAGCGTGATT GGAAGACAGC TGTACACGTC 501 AAGTGAGCGG GAAGATGCAG CAGCGTGATT GGAAGACAGC TGTACACGTC 501 AAGTGAGCGG GAAGATGCAG CAGCGTGATT GGAAGACAGC TGTACACGTC	550 550 550

FIG. 3a

600 600 600



CLASS SUBCLASS

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APPROVED O.G. FIG.

Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

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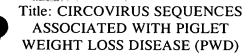
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	<u>610</u> <u>620</u> <u>630</u> <u>640</u> <u>650</u>	
circopormank circopormeeh circopordfp	601 TGAGCCTAGC GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG 601 TGAGCCTAGG GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG 601 TGAGCCTAGG GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG	650 650 650
circopormank circopormeeh circopordfp	660 670 680 690 700 651 GATATCATGG AGAAGAAGTT GTTGTTTTGG ATGATTTTTA TGACTGGTTA 651 GATATCATGG AGAAGAAGTT GTTGTTTTGG ATGATTTTTA TGACTGGTTA 651 GATATCATGG AGAAGAAGTT GTTGTTTTGG ATGATTTTTA TGACTGGTTA	700 700 700
circopormank circopormeeh circopordfp	710 720 730 740 750 701 CCTTGGGATG ATCTACTGAG ACTGTGTGAC CGGTATCCAT TGACTGTAGA	750 750 750
circopormank circopormeeh circopordfp	760 770 780 790 800 751 GACTAAAGGC GGTACTGTTC CTTTTTTGGC TCGCAGTATT TTGATTACCA 751 GACTAAAGGG GGTACTGTTC CTTTTTTGGC CCGCAGTATT TTGATTACCA 751 GACTAAAGGG GGTACTGTTC CTTTTTTGGC CCGCAGTATT TTGATTACCA	800 800 800
circopormank circopormeeh- circópordfp	810 820 830 840 850 801 GCAATCAGGC CCCCCAGGAA TGGTACTCCT CAACTGCTGT CCCAGCTGTA 801 GCAATCAGGC CCCCCAGGAA TGGTACTCCT CAACTGCTGT CCCAGCTGTA 801 GCAATCAGGC CCCCCAGGAA TGGTACTCCT CAACTGCTGT CCCAGCTGTA	850 850 850
circopormank circopormeeh circopordfp	860 870 880 890 900 851 GAAGCTCTCT ATCGGAGGAT TACTACTTTG CAATTTTGGA AGACTGCTGG 851 GAAGCTCTCT ATCGGAGGAT TACTACTTTG CAATTTTGGA AGACTGCTGG 851 GAAGCTCTCT ATCGGAGGAT TACTACTTTG CAATTTTGGA AGACTGCTGG	900 900 900
circopormank circopormeeh circopordfp	910 920 930 940 950 901 AGAACAATGA ACGGAGGTAC CCGAAGGCCG ATTTGAAGCA GTGGACCCAC 901 AGAACAATCC ACGGAGGTAC CCGAAGGCCG ATTTGAAGCA GTGGACCCAC 901 AGAACAATCC ACGGAGGTAC CCGAAGGCCG ATTTGAAGCA GTGGACCCAC	950 950 950
circopormank circopormeeh circopordfp	960 970 980 990 1000 951 CCTGTGCCCT TTTCCCATAT AAAATAAATT ACTGAGTCTT TTTTGTTATC 951 CCTGTGCCCT TTTCCCATAT AAAATAAATT ACTGAGTCTT TTTTGTTATC 951 CCTGTGCCCT TTTCCCATAT AAAATAAATT ACTGAGTCTT TTTTGTTATC	1000 1000 1000
circopormank circopormeeh circopordfp	1010 1020 1030 1040 1050 1001 ACATCGTAATI GGTTTTTATTI TTTATTTATTI TAGAGGGTCTI TTTAGGATAA 1001 ACATCGTAATI GGTTTTTATTI TTTATTI TAGAGGGTCTI TTTAGGATAA 1001 ACATCGTAATI GGTTTTTATTI TTTATTICATTI TAGAGGGTCTI TTCAGGATAA	1050 1050 1050
circopormank circopormeeh circopordfp	1060 1070 1080 1090 1100 1051 ATTCTCTGAA TTGTACATAA ATAGTCAGCC TTACCACATA ATTTTGGGCT 1051 ATTCTCTGAA TTGTACATAA ATAGTCAGCC TTACCACATA ATTTTGGGCT 1051 ATTCTCTGAA TTGTACATAA ATAGTCAACC TTACCACATA ATTTTGGGCT	1100 1100 1100
circopormank circopormeeh circopordfp	1110 1120 1130 1140 1150 1101 GTGGCTGCAT TTTGGAGCGC ATAGCCGAGG CCTGTGTGCT CGACATTGGT 1101 GTGGCTGCAT TTTGGAGCGC ATAGCCGAGG CCTGTGTGCT CGACATTGGT 1101 GTGGTTGCAT TTTGGAGCGC ATAGCCGAGG CCTGTGTGCT CGACATTGGT	1150 1150 1150
circopormank circopormeeh circopordfp	1160 1170 1180 1190 1200 1151 GTGGGTATTT AAATGGAGCC ACAGCTGGTT TCTTTTATTA TTTTGGTTGGA 1151 GTGGGTATTT AAATGGAGCC ACAGCTGGTT TCTTTTATTA TTTTGGTTGGA 1151 GTGGGTATTT AAATGGAGCC ACAGCTGGTT TCTTTTATTA TTTTGGTTGGA	1200 1200 1200



APPROVED O.G. FIG.

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Inventors: Jestin et al. Appl. No.: 09/514,245

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	1210	1220 1230		
-circopormank -circopormeeh	1201 ACCAATCAAT TGT	TTGGTCC AGCTCAGGTT	T TGGGGGTGAA GTACCTGGAG T TGGGGGTGAA GTACCTGGAG T TGGGGGTGAA GTACCTGGAG	1250 1250 1250
circopormank circopormeeh circopordfp	1260 1251 <mark>TGGTAGGTAA AGG</mark> 1251 TGGTAGGTAA AGG 1251 <u>TGGTAGGTAA AGG</u>	GCTGCCT TATGGTGTGC	1290 CGGGAGGAGT AGTTAATATA CGGGAGGAGT AGTTAATATA CGGGAGGAGT AGTTAATATA	1300 1300 1300
circopormank circopormeeh circopordfp	1301 GGGGTCATAG IGCC	AAGTTGGIITGGAGGGGGT	1340 TTACAAAGTTG GCATCCAAGA TTACAAAGTTG GCATCCAAGA TTACAAAGTTG GCATCCAAGA	1350 1350 1350
circopormank circopormeeh circopordfp	1351 ITAACAACAGTI IGGA	CCCAACA ICCTCTTTGAT	1390 1400 TITAGAGGTGATI GGGGTCTCTG TAGAGGTGATI GGGGTCTCTG TAGAGGTGATI GGGGTCTCTG	1400 1400 1400
circopormank circopormeeh circopordfp	1401 GGGTAAAATTI CAT	ATTTAGC CTTTCTAATA	1440 1 CGGTAGTATT GGAAAGGTAG 1 CGGTAGTATT GGAAAGGTAG 1 CGGTAGTATT GGAAAGGTAG	1450 1450 1450
circopormank circopormeeh circopordfp	1451 GGGTAGGGGG TTG	GTGCCGC CTGAGGGGGG	1490 GAGGAACTGG CCGATGTTGA GAGGAACTGG CCGATGTTGA GAGGAACTGG CCGATGTTGA	1500 1500 1500
circopormank circopormeeh circopordfp	1501 ATTTGAGGTA GTT	1520 AACATGC CAAGATGGCT AACATTC CAAGATGGCT AACATTC CAAGATGGCT	GCGAGTATCC TCCTTTTATG GCGAGTATCC TCCTTTTATG	1550 1550 1550
circopormank circopormeeh circopordfp	1551 GTGAGTACAA ATT	CTGTAGA AAGGCGGGAA	1590 TTGAAGATAC CCGTCTTTCG TTGAAGATAC CCGTCTTTCG TTGAAGATAC CCGTCTTTCG	1600 1600 1600
circopormank circopormeeh circopordfp	1601 [GCGCCATCTG] [TAA	CGGTTTC TGAAGGCGGG	1640 GTGTGCCAAA TATGGTCTTC GTGTGCCAAA TATGGTCTTC GTGTACCAAA TATGGTCTTC	1650 1650 1650
circopormank circopormeeh circopordfp	1660 1651 TCCGGAGGAT GTT 1651 TCCGGAGGAT GTT 1651 TCCGGAGGAT GTT	TCCAAGA ITGGCTGCGGG	1690 GGCGGGTCCT TCTTCTGCGG GGCGGGTCCT TCTTCTGCGG GGCGGGTCCG TCTTCTGCGG	1700 1700 1700
circopormank circopormeeh circopordfp	1701 TAACGCCTCC TTG	GCCACGT CATCCTATAA	1740 AAGTGAAAGA AGTGCGCTGC AAGTGAAAGA AGTGCGCTGC AAGTGAAAGA AGTGCGCTGC	1750 1750 1750
circopormank circopormeeh circopordfp	1760 1751 TGTAGTATT 1751 TGTAGTATT 1751 TGTAGTATT	1770 1780		1800 1800 1800



APPROVED O.G. FIG.
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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

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	10 20 30 40 50	
circopormank circopormeeh	1 NPSKKSGPOP HKRWVFTLNN PSEEEKNKIR ELPISLFDYF VCGEEGLEEG 1 NPSKKSGPOP HKRWVFTLNN PSEEEKNKIR ELPISLFDYF VCGEEGLEEG	50 50 50
circopordfp[1 NPSKKSGPOP HKRWVFTLNN PSEEEKNKIR ELPISLEDYF VCGEEGLEEG	5ŏ
circopormank	60 70 80 90 100 51 RTAHLOGFAN FAKKOTFNKV KWYFGARCHI EKAKGTDOON KEYCSKEGHI	100
circopormeeh	51 RTPHLÖGFAN FAKKÖTFNKV KWYFGARCHI EKAKGTDÖÖN KEYCSKEGHI	$\begin{array}{c} 100 \\ 100 \end{array}$
circopordfp[51 RTPHLÖGFAN FAKKÖTFNKV KWYFGARCHI EKAKGTDOON KEYCSKEGHI	100
circopormank	110 120 130 140 150 101 LIECGAPRNO GKRSDLSTAV STLLETGSLV TVAEQFPVTY VRNFRGLAEL	150
circopormeeh	101[LIECGAPRNO [GKRSDLSTAV][STLLETGSLV][TVAEQFPVTY][VRNFRGLAEL]	150-
circopordfp[101 LIECGAPRNO GKRSDLSTAV STLLETGSLV TVAEOFPVTY VRNFRGLAEL	150
circopormank	160 170 180 190 200 151[EKVSGKMOOR][DWKTAVHVIV][GPPGCGKSOW][ARNFAEPSOT][YWKPSRNKWH]	200
circopormeeh	151 LKVSGKMÕÕR DWKTAVHVIV GPPGCGKSÕW ARNFAEPROT YWKPSRNKWH	200
circopordfp[151 LKVSGKMQQR DWKTAVHVIV GPPGCGKSQW ARNFAEDROT YWKDSRNKWH	200
circopormank '	210 220 230 240 250 201 DGYHGEEVVV EDDFYGWLPW DDLLRLCDRY PLTVETKGGT VPFLARSILI	250
circopormeeh	201 DGYHGEEVVV LDDFYGWLPW DDLLRLCDRY PLTVETKGGT VPFLARSILI	250 250
circopordfp[201 DGYHGEEVVV LDDFYGWLPW DDLLRLCDRY PLTVETKGGT VPFLARSILI	250
; circopormank	260 - 270 280 290 300 251 TSNQAPQEWY SSTAVPAVEA LYRRITTLQF WKTAGEQSTE VPEGRFEAVD	300
circopormeeh	251 TSNQAPQEWY SSTAVPAVEA LYRRITTLQF WKTAGEQSTE VPEGRFEAVD	300
circopordfp[251 TSNQAPQEWY SSTAVPAVEA LYRRITTLQF WKTAGEQSTE VPEGRFEAVD	300
circopormank	310 320 330 340 350	250
circopormeeh	301 PPCALFPYKI NY	350 350 350
circopordfp[301 PPCALFPYKI NY	350

FIG. 4



APPROVED O.G. FIG.

BY CLASS SUBCLASS

Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLE WEIGHT LOSS DISEASE (PW) Inventors: Jestin et al. Appl. No.: 09/514,245

circopormank circopormeeh circopordfp[10 20 30 40 50 1 MTWPRRYRR RRTRPRSHLG NILRRPYLA HPAFRNRYRW RRKTGIFNCR 1 MTWPRRYRR RRTRPRSHLG NILRRPYLA HPAFRNRYRW RRKTGIFNSR 1 MTWPRRYRR RRTRPRSHLG NILRRPYLV HPAFRNRYRW RRKTGIFNSR	50 50 50
circopormank circopormeeh circopordfp[60 70 100 51 LSKEFVITIK GGYSOPSWIM NILRFNIGOF LPPSGGTNPL PLPFOYYRIR 51 LSTEFVLTIK GGYSOPSWNV NYLKFNIGOF LPPSGGTNPL PLPFOYYRIR 51 LSREFVLTIR GGISOPSWNV NELRFNIGOF LPPSGGTNPL PLPFOYYRIR	100 100 100
circopormank circopormeeh circopordfp[110 120 130 140 150 101 KAKYEFYPRD PITSNERGYG STVVILDANF VTPSTNLAYD PYINYSSRHT 101 KAKYEFYPRD PITSNORGYG STVVILDANF VTPSTNLAYD PYINYSSRHT 101 KAKYEFYPRD PITSNORGYG STVVILDANF VTPSTNLAYD PYINYSSRHT	150 150 150
circopormank circopormeeh circopordfp[160 170 180 190 200 151 TROPFTYHSR YFTPKPELDO TIEWFHPNNK RNOLWLHLNT HTNVEHTGLG 151 TROPFTYHSR YFTPKPELDO TIEWFHPNNK RNOLWLHLNT HTNVEHTGLG 151 TROPFTYHSR YFTPKPELDO TIEWFOPNNK RNOLWLHLNT HTNVEHTGLG	200 200 200
circopormank circopormeeh circopordfp[210 220 230 240 250 201 YALQNAATAQ NYVVRLTIYV QFREFILKDP LNK* 201 YALQNAATAQ NYVVRLTIYV QFREFILKDP LNK* 201 YALQNATTAQ NYVVRLTIYV QFREFILKDP LNE*	250 250 250

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circopormank circopormeeh circopordfp[1 1 1		20 RLPVGVARLS RLPVGVPRLS RLPVGVPRLS	KITGPLALPT	TGRAHYDVYS	CLPITLLHLP CLPITLLHLP	50 50 50
circopormank circopormeeh circopordfp[51 51 51	AHFÕKFSÕPA	70 EISHIRYREL EISHIRYREL EISHIRYRKL	LGYSHÔRPRL	QKGTHSSRQV	100 AALPLVPRSS AALPLVPRSS AALPLVPRSS	100 100 100
circopormank circopormeeh circopordfp[101 101 101	110 TLDKYVAFFT TLDKYVAFFT TLDKYVAFFT	AVFFILLVGS			150 LUSKISKPLE LUSKIRKPLE LUSKIRKPLE	150 150 150
circopormank circopormeeh circopordfp[151 151 151	160 MSSSTLFOTF VRSSTLFOTF VRSSTLFOTF	170 LSANKIIKKG LSANKIIKKG LATNKIIKKG	180 DWKLPYFVFL DWKLPYFVFL DWKLPYFVFL	LLGRIIKGEH	200 PPLMGLRAAF PPLMGLRAAF PPLMGLRAAF	200 200 200
circopormank circopormeeh circopordfp[201 201 201	210 LAWHFH* LAWHFEH LAWHFH	220	230	240	250	250 250 250

FIG. 6



Title: CIRCOVIRUS SEQUENCES
SOCIATED WITH PIGLET
WEIGHT LOSS DISEASE (PWD)
Inventors: Jestin et al.

Appl. No.: 09/514,245

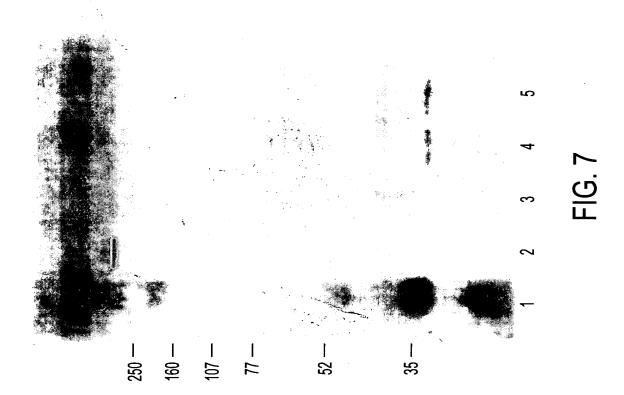
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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

> Inventors: Jestin et al. Appl. No.: 09/514,245

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0 ₽ Leu Ala Ser Arg Cys Arg Cys Arg Pro Leu Val Glu Ala Ala Val His Gly Trp Arg Val Glu Ala Ala Ala Ala Gly Arg Cys Cys Arg Leu Leu Met Gly Gly Ala Cys Lys Pro Leu Pro Leu Val Glu Ala Ala Gly *** Cys Cys Ala TGG TCG CGT GAA GCC GTC GCC GTC GTG GAG CCG TCG TGG AGT CGT CGT TGT ACG 27 ACC AGC GCA CTT CGG CAG CGG CAC CTC GGC AGC ACC TCA GCA GCA ACA TGC Thr Ser Ala Leu Arg Gln Arg Gln His Leu Gly Ser Thr Ser Ala Ala Thr Cys Pro Ala His Phe Gly Ser Gly Ser Thr Ser Ala Ala Pro Gln Gln His Ala GIn Arg Thr Ser Ala Ala Ala Ala Pro Arg Gln His Leu Ser Ser Asn Met Pro

Ala Leu Leu Ile Ser Ser Ala Ser Gly Leu Gly Met Phe Pro Pro His Glu Ser Leu Leu Phe Phe Pro Leu Leu Pro Gly Trp Gly Trp Leu Leu His Thr Asn Val Trp Cys Ser Ser His Phe Phe Arg Val Gly Val Gly Tyr Phe Thr Pro Thr *** TCT TAC CTT CTT CGC CTG GGG TTG GGG TAT TTT CCA CCC ACA AGT 72 81 90 99 108 GGT CGT TCT 108 CCA GCA AGA AGA ATG GAA GAA GCG GAC CCC AAC CCC ATA AAA GGT GGG TGT TCA Pro Ala Arg Arg Met Glu Glu Ala Asp Pro Asn Pro Ile Lys Gly Gly Cys Ser Gln Gln Glu Glu Trp Lys Lys Arg Thr Pro Thr Pro *** Lys Val Gly Val His Ser Lys Lys Asn Gly Arg Ser Gly Pro Gln Pro His Lys Arg Trp Val Phe Thr

Gln Ilè Ile Arg Gly Phe Val Leu Ala Leu Phe Tyr Pro Ile Lys Trp Tyr Gly Arg Phe Leu Gly Glu Ser Ser Ser Arg Leu Phe Ile Arg Ser Arg Gly Ile Asp Glu Ser Tyr Asp Lys Arg Leu Arg Ala Cys Ser Phe Val Pro Asp Glu Leu Ile GAG ACT TAT TAG GAA GGC TTC TGC TCG CGT TCT TTT ATG CCC TAG AAG GTT ATA 117 126 135 144 153 162 CTC TGA ATA ATC CTT CCG AAG ACG AGC GCA AGA AAA TAC GGG ATC TTC CAA TAT Leu *** Ile Ile Leu Pro Lys Thr Ser Ala Arg Lys Tyr Gly Ile Phe Gln Tyr Ser Glu *** Ser Phe Arg Arg Arg Ala Gln Glu Asn Thr Gly Ser Ser Asn Ile Leu Asn Asn Pro Ser Glu Asp Glu Arg Lys Lys Ile Arg Asp Leu Pro Ile Ser

*** Lys Ile Ile Lys Asn Asn Ala Leu Leu Thr Ile Leu Phe Ser Ser Cys Arg Arg Asn Ser *** Lys Ile Thr Pro Ser Ser Pro Leu Ser Ser Pro Arg Val Gly Ile Gln Asn Asn *** Gln Gln Arg Pro Pro Tyr His Pro Leu Val Phe Val GGG ATA AAC TAA TAA AAT AAC AAC CGC TCC TCC CAT TAC TCC TTC CTG CTT GTG 171 180 189 198 207 216 CCC TẠT TTG ATT ATT TTA TTG TTG GCG AGG AGG GTA ATG AGG ĀAG GAC GAA CÁC Pro Tyr Leu Ile Ile Leu Leu Leu Ala Arg Arg Val Met Arg Lys Asp Glu His Pro Ile *** Leu Phe Tyr Cys Trp Arg Gly Gly *** *** Gly Arg Thr Asn Thr Leu Phe Asp Tyr Phe Ile Val Gly Glu Glu Gly Asn Glu Glu Gly Arg Thr Pro

Val Glu Leu Pro Glu Ser Ile Lys His Leu Leu Leu Ser Lys Ile Phe His Leu
*** Arg Trp Pro Asn Ala Leu Lys Thr Phe Phe Cys Val Lys Leu Leu Thr Phe
Glu Gly Gly Pro Thr Arg *** Asn Gln Ser Ser Ala Ser Lys *** Tyr Leu Ser GAG TGG AGG TCC CCA AGC GAT TAA AAC ACT TCT TCG TCT GAA AAT TAT TTC ACT 261 CTC ACC TCC AGG GGT TCG CTA ATT TTG TGA AGA ĀGC AGA CTT TTĀ ATA AAG TGA Leu Thr Ser Arg Gly Ser Leu Ile Leu *** Arg Ser Arg Leu Leu Ile Lys ***
Ser Pro Pro Gly Val Arg *** Phe Cys Glu Glu Ala Asp Phe *** *** Ser Glu
His Leu Gln Gly Phe Ala Asn Phe Val Lys Lys Gln Thr Phe Asn Lys Val Lys



SUBCLASE

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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD) Inventors: Jestin et al.

Inventors: Jestin et al. Appl. No.: 09/514,245

Pro Ile Gln Thr Gly Ala Ala Val Asp Leu Phe Arg Phe Ser Cys Ile Leu Leu His Tyr Lys Pro Ala Arg Gln Trp Met Ser Phe Ala Phe Pro Val Ser *** Cys Thr Thr Asn Pro His Gly Ser Gly Cys Arg Ser Leu Ser Leu Phe Leu Asp Ala TCA CCA TAA ACC CAC GGG CGA CGG TGT AGC TCT TTC GCT TTC CTT GTC TAG TCG 279 288 297 306 315 324 AGT GGT ATT TGG GTG CCC GCT GCC ACA TCG AGA AAG CGA AAG GAA CAG ATC AGC Ser Gly Ile Trp Val Pro Ala Ala Thr Ser Arg Lys Arg Lys Glu Gln Ile Ser Val Val Phe Gly Cys Pro Leu Pro His Arg Glu Ser Glu Arg Asn Arg Ser Ala Trp Tyr Leu Gly Ala Arg Cys His Ile Glu Lys Ala Lys Gly Thr Asp Gln Gln

Ile Phe Phe Val Ala Thr Phe Phe Ala Val *** Gln His Leu Thr Ser Ser Arg Phe Leu Ser Tyr Gln Leu Leu Ser Pro Leu Lys Ser His Ser His Pro Ala Gly Ser Tyr Leu Ile Ser Cys Tyr Leu Leu Cys Ser Val Ser Pro Thr His Leu Glu TCT TAT TTC TTA TGA CGT CAT TTC TTC CGT TGA ATG ACT ACC TCA CAC CTC GAG 333 342 351 360 369 378 AGA ATA AAG AAT ACT GCA GTA AAG AAG GCA ACT TAC TGA TGG AGT GTG GAG CTC Arg Ile Lys Asn Thr Ala Val Lys Lys Ala Thr Tyr *** Trp Ser Val Glu Leu Glu *** Arg Ile Leu Gln *** Arg Arg Gln Leu Thr Asp Gly Val Trp Ser Ser Asn Lys Glu Tyr Cys Ser Lys Glu Gly Asn Leu Leu Met Glu Cys Gly Ala Pro

Ser Arg Leu Ser Leu Pro Thr Val Gln Arg Ser Ser His Thr Gly Gln Gln Leu Leu Asp *** Pro Cys Arg Leu Ser Arg Asp Val Ala Thr Leu Val Lys Asn Ser *** Ile Glu Pro Val Val Ser His Gly Thr *** Gln Gln Ser Tyr Arg Thr Pro GAT CTA GAG TCC CTG TTG CCT CAC TGG ACA GAT GAC GAC ACT CAT GGA ACA ACC 387 396 405 414 423 432 432 CTA GAT CTC AGG GAC AAC GGA GTG ACC TGT CTA CTG CTG TGA GTA CCT TGT TGG Leu Asp Leu Arg Asp Asn Gly Val Thr Cys Leu Leu Leu *** Val Pro Cys Trp *** Ile Ser Gly Thr Thr Glu *** Pro Val Tyr Cys Cys Glu Tyr Leu Val Gly Arg Ser Gln Gly Gln Arg Ser Asp Leu Ser Thr Ala Val Ser Thr Leu Leu Glu

Ala Pro Thr Gln His Gly Asn Cys Leu Leu Val Arg Tyr Arg Lys Asp Ser Ile Leu Pro Leu Arg Thr Val Thr Ala Ser Cys Cys Gly Thr Val Asn Thr Leu Phe Ser Arg Ser Asp Pro Ser Arg Gln Leu Ala Ala Gly Gln Leu Thr Gln *** Phe TCT CGC CCT CAG ACC ACT GGC AAC GTC TCG TCG TGG GAC ATT GCA AAC AGT CTT 441 450 459 468 477 486 AGA GCG GGA GTC TGG TGA CCG TTG CAG AGC ACC CTG TAA CGT TTG TCA GAA Arg Ala Gly Val Trp *** Pro Leu Gln Ser Ser Thr Leu *** Arg Leu Ser Glu Glu Arg Glu Ser Gly Asp Arg Cys Arg Ala Ala Pro Cys Asn Val Cys Gln Lys Ser Gly Ser Leu Val Thr Val Ala Glu Gln His Pro Val Thr Phe Val Arg Asn

Glu Ala Pro Gln Ser Phe Lys Gln Phe His Ala Pro Phe His Leu Leu Thr Ile Lys Arg Pro Ser Ala Ser Ser Lys Phe Thr Leu Pro Phe Ile Cys Phe Arg Ser Asn Gly Arg Ala Pro Gln Val Lys Ser Leu Ser Arg Ser Phe Ala Ser Ala His TAA AGG CGC CCG ACC GAC TTG AAA ACT TTC ACT CGC CCT TTT ACG TCT TCG CAC 495 504 513 522 531 540 ATT TCC GCG GGC TGG CTG AAC TTT TGA AAG TGA GCG GGA AAA TGC AGA AGC GTG Ile Ser Ala Gly Trp Leu Asn Phe *** Lys *** Ala Gly Lys Cys Arg Ser Val Phe Pro Arg Ala Gly *** Thr Phe Glu Ser Glu Arg Glu Asn Ala Glu Ala *** Phe Arg Gly Leu Ala Glu Leu Leu Lys Val Ser Gly Lys Met Gln Lys Arg Asp



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FIG.

Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

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Pro Leu Ser Ile Tyr Val Asp Asn His Pro Trp Arg Pro Thr Thr Phe Ala Phe Gln Phe Val Leu Thr Cys Thr Met Thr Pro Gly Gly Pro His Pro Leu Leu Leu Asn Ser Ser *** His Val Arg *** Gln Pro Ala Val Gln Thr His Tyr Phe Cys

TAA CCT TCT GAT TAC ATG TGC AGT AAC ACC CCG GTG GAC CCA CAC CAT TTT CGT 549 558 567 576 585 594

ATT GGA AGA CTA ATG TAC ACG TCA TTG TGG GGC CAC CTG GGT GTG GTA AAA GCA

Ile Gly Arg Leu Met Tyr Thr Ser Leu Trp Gly His Leu Gly Val Val Lys Ala Leu Glu Asp *** Cys Thr Arg His Cys Gly Ala Thr Trp Val Trp *** Lys Gln Trp Lys Thr Asn Val His Val Ile Val Gly Pro Pro Gly Cys Gly Lys Ser Lys

Pro Ser Ser Ile Lys Cys Val Arg Phe Gly Cys Val Pro Phe Trp Arg Ser Val His Ala Ala Leu Lys Ala Ser Gly Ser Val Val Tyr Gln Phe Gly Gly Leu Phe Ile Pro Gln *** Asn Gln Leu Gly Pro Phe Trp Met Ser Ser Val Val *** Phe

TTA CCC GAC GAT TAA AAC GTC TGG GCC TTT GGT GTA TGA CCT TTG GTG GAT CTT
603 612 621 630 639 648

AAT GGG CTG CTA ATT TTG CAG ACC CGG AAA CCA CAT ACT GGA AAC CAC CTA GAA

Asn Gly Leu Leu Ile Leu Gln Thr Arg Lys Pro His Thr Gly Asn His Leu Glu
Met Gly Cys *** Phe Cys Arg Pro Gly Asn His Ile Leu Glu Thr Thr *** Lys
Trp Ala Ala Asn Phe Ala Asp Pro Glu Thr Thr Tyr Trp Lys Pro Pro Arg Asn

Leu Pro Pro Ile Thr Val Met Thr Phe Phe His Asn Asn Asn Ile Val Lys Ile Leu His His Ser Pro *** Trp Pro Ser Ser Thr Thr Thr Ile Ser Ser Lys *** Cys Thr Thr Pro His Asn Gly His His Leu Leu Pro Gln *** Gln His Ser Lys TGT TCA CCA CCC TAC CAA TGG TAC CAC TTC TTC ACC AAC AAT AAC TAC TGA AAA 657 666 675 684 693 702 ACA AGT GGT GGG ATG GTT ACC ATG GTG AAG AAG TGG TTG TTA TTG ATG ACT TTT Thr Ser Gly Gly Met Val Thr Met Val Lys Lys Trp Leu Leu Leu Met Thr Phe Gln Val Val Gly Trp Leu Pro Trp *** Arg Ser Gly Cys Tyr *** *** Leu Leu Lys Trp Trp Asp Gly Tyr His Gly Glu Glu Val Val Val Ile Asp Asp Phe Tyr

Ala Pro Gln Gly Pro Ile Ile *** Gln Ser Gln Thr Ile Ser Ile Trp Gln Ser Pro Gln Ser Gly Gln Ser Ser Arg Ser Leu Ser His Ser Arg Tyr Gly Asn Val His Ser Ala Ala Arg Pro His Asp Val Ser Val Thr His Asp Ile Asp Met Ser

TAC CGA CCG ACG GGA CCC TAC TAG ATG ACT CTG ACA CAC TAG CTA TAG GTA ACT
711 720 738 747 756

ATG GCT GGC TGC CCT GGG ATG ATC TAC TGA GAC TGT GTG ATC GAT ATC CAT TGA

Met Ala Gly Cys Pro Gly Met Ile Tyr *** Asp Cys Val Ile Asp Ile His ***

Trp Leu Ala Ala Leu Gly *** Ser Thr Glu Thr Val *** Ser Ile Ser Ile Asp
Gly Trp Leu Pro Trp Asp Asp Leu Leu Arg Leu Cys Asp Arg Tyr Pro Leu Thr

Tyr Leu Ser Phe Thr Ser Ser Tyr Arg Lys Gln Gly Ala Thr Asn Gln Asn Gly Thr Ser Val Leu Pro Pro Val Thr Gly Lys Lys Ala Arg Leu Ile Arg Ile Val Gln Leu Ser *** Leu His Phe Gln Val Lys Lys Pro Gly Cys Tyr Glu Ser ***

GAC ATC TCT GAT TTC CAC CTT GAC ATG GAA AAA ACC GGG CGT CAT AAG ACT AAT 765 774 783 792 801 810

CTG TAG AGA CTA AAG GTG GAA CTG TAC CTT TTT TGG CCC GCA GTA TTC TGA TTA Leu *** Arg Leu Lys Val Glu Leu Tyr Leu Phe Trp Pro Ala Val Phe *** Leu Cys Arg Asp *** Arg Trp Asn Cys Thr Phe Phe Gly Pro Gln Tyr Ser Asp Tyr Val Glu Thr Lys Gly Gly Thr Val Pro Phe Leu Ala Arg Ser Ile Leu Ile Thr

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FIG.

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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

Ala Ile Leu Gly Arg Gln Phe Pro Val Gly *** Ser Ser Asp Trp Ser Tyr Phe Leu Leu *** Val Gly Asn Ser His Tyr Glu Glu Val Ala Thr Gly Ala Thr Ser Trp Cys Asp Ser Gly Thr Pro Ile Thr Ser Arg Leu Gln Gln Gly Leu Gln Leu GGT-CGT-TAG-TCT-GGG-GCA-ACC-TTA-CCA-TGA-GGA-GTT-GAC-GAC-AGG-GTC-GAC-ATC-819 828 837 846 855 864 CCA GCA ATC AGA CCC CGT TGG AAT GGT ACT CCT CAA CTG CTG TCC CAG CTG TAG-Pro Ala Ile Arg Pro Arg Trp Asn Gly Thr Pro Gln Leu Leu Ser Gln Leu *** Gln Gln Ser Asp Pro Val Gly Met Val Leu Leu Asn Cys Cys Pro Ser Cys Arg Ser Asn Gln Thr Pro Leu Glu Trp Tyr Ser Ser Thr Ala Val Pro Ala Val Glu

Ser Lys Ile Pro Pro Asn Ser Gly Gln Tyr Lys Pro Leu Ile Ser Cys Phe Leu Ala Arg *** Arg Leu Ile Val Glu Lys Thr Asn Gln Phe Phe Ala Val Ser Cys Leu Glu Lys Asp Ser Ser *** Lys Arg Pro Ile Lys Ser Ser His *** Leu Val TTC GAG AAA TAG CCT CCT AAT GAA GGA ACC ATA AAA CCT TCT TAC GAT GTC TTG 873 882 891 900 909 918 AAG CTC TTT ATC GGA GGA TTA CTT CCT TGG TAT TTT GGA AGA ATG CTA CAG AAC Lys Leu Phe Ile Gly Gly Leu Leu Pro Trp Tyr Phe Gly Arg Met Leu Gln Asn Ser Ser Leu Ser Glu Asp Tyr Phe Leu Gly Ile Leu Glu Glu Cys Tyr Arg Thr Ala Leu Tyr Arg Arg Ile Thr Ser Leu Val Phe Trp Lys Asn Ala Thr Glu Gln

Gly Arg Leu Phe Pro Ala Leu Glu Asp Gly Lys Gly Gly Trp Ala Arg Phe Lys Asp Val Ser Ser Pro Pro Trp Asn Thr Val Arg Glu Gly Gly His Gly Ser Asn Ile Trp Pro Pro Leu Pro Gly Thr Arg *** Gly Lys Gly Gly Met Gly Gln Ile

TTA GGT GCC TCC TTC CCC CGG TCA AGC AGT GGG AAA GGG GGG GTA CGG GAC TTA 927

AAT CCA CGG AGG AAG GGG GCC AGT TCG TCA CCC TTT CCC CCC CAT GCC CTG AAT

Asn Pro Arg Arg Lys Gly Ala Ser Ser Ser Pro Phe Pro Pro His Ala Leu Asn Ile His Gly Gly Arg Gly Pro Val Arg His Pro Phe Pro Pro Met Pro *** Ile Ser Thr Glu Glu Gly Gly Gln Phe Val Thr Leu Ser Pro Pro Cys Pro Glu Phe

Trp Ile Phe Tyr Ile Val Ser Asp Lys Lys Asp Ser Arg Leu Pro Lys *** *** Gly Tyr Ser Ile Phe *** Gln Thr Lys Lys Ile Val Glu Tyr His Asn Lys Asn Glu Met His Phe Leu Asn Ser Leu Arg Lys *** *** Lys Thr Ile Thr Lys Ile

AAG GTA TAC TTT ATT TAA TGA CTC AGA AAA AAT AGT GAA GCA TTA CCA AAA ATA 981 990 1008 1017 1026

TTC CAT ATG AAA TAA ATT ACT GAG TCT TTT TTA TCA CTT CGT AAT GGT TTT TAT

Phe His Met Lys *** Ile Thr Glu Ser Phe Leu Ser Leu Arg Asn Gly Phe Tyr Ser Ile *** Asn Lys Leu Leu Ser Leu Phe Tyr His Phe Val Met Val Phe Ile Pro Tyr Glu Ile Asn Tyr *** Val Phe Phe Ile Thr Ser *** Trp Phe Leu Leu Leu

Glu Asn Leu Thr Leu His Pro Thr Lys Leu Ile Leu Asn Glu Ser Asn Tyr Met Asn Met Leu Pro *** Thr Pro Pro Arg *** Phe *** Ile Arg Gln Ile Thr Cys Ile *** *** Pro Asn Leu Pro Pro Asp Lys Phe Asn Phe Glu Arg Phe Gln Val ATA AGT AAT TCC CAA TTC ACC CCC CAG AAA TTT TAA TTT AAG AGA CTT AAC ATG 1035 1044 1053 1062 1071 1080 TAT TCA TTA AGG GTT AAG TGG GGG GTC TTT AAA ATT AAA TTC TCT GAA TTG TAC Tyr Ser Leu Arg Val Lys Trp Gly Val Phe Lys Ile Lys Phe Ser Glu Leu Tyr Ile His *** Gly Leu Ser Gly Gly Ser Leu Lys Leu Asn Ser Leu Asn Cys Thr Phe Ile Lys Gly *** Val Gly Gly Leu *** Asn *** Ile Leu *** Ile Val His



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ASSOCIATED WITH PIGLET EIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

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Cys Pro *** Val Ser Ile Thr Asn Arg Thr Thr Tyr Val Thr Lys Ser Arg Leu Val His Asn Cys Pro Tyr Gln Ile Gly Pro Arg Ile Tyr Gln Lys Arg Val Cys Tyr Met Thr Val Arg Ile Asn Tyr Glu Gln Asp Tyr Ile Ser Asn Glu Phe Ala TAT GTA CCA ATG TGC CTA TAA CAT AAG GAC CAG CAT ATA TGA CAA AAG CTT GCG 1089 1098 1107 1116 1125 1134 ATA CAT GGT TAC ACG GAT ATT GTA TTC CTG GTC GTA TAT ACT GTT TTC GAA CGC Ile His Gly Tyr Thr Asp Ile Val Phe Leu Val Val Tyr Thr Val Phe Glu Arg Tyr Met Val Thr Arg Ile Leu Tyr Ser Tyr Ile Leu Phe Ser Asn Ala Thr Trp Leu His Gly Tyr Cys Ile Pro Gly Arg Ile Tyr Cys Phe Arg Thr Gln

Ala Ser Ala *** Thr Thr *** Met Glu Leu Leu Lys Tyr Asp *** Gly Cys Ser His Arg Pro Arg Arg Pro Arg Cys Lys Trp Cys Asn Thr Thr Glu Ala Val Ala Thr Gly Leu Gly Val His Asp Val Asn Gly Ala Thr Gln Leu Arg Leu Trp Leu TCA CGG CTC CGG ATG CAC CAG ATG TAA AGG TCG TCA AAC ATC AGA GTC GGT GTC 1143 1152 1161 1170 1179 1188 AGT GCC GAG GCC TAC GTG GTC TAC ATT TCC AGC AGT TTG TAG TCT CAG CCA CAG Ser Ala Glu Ala Tyr Val Val Tyr Ile Ser Ser Ser Leu *** Ser Gln Pro Gln Val Pro Arg Pro Thr Trp Ser Thr Phe Pro Ala Val Cys Ser Leu Ser His Ser Cys Arg Gly Leu Arg Gly Leu His Phe Gln Gln Phe Val Val Ser Ala Thr Ala

Pro Pro Leu Thr Gly Pro Thr Thr Pro Ser Pro Ser Pro *** Pro Ile Ala Pro Gln Pro Tyr Leu Val Pro Leu Pro Leu Leu Leu Ala Pro Asn His Tyr Pro Pro Lys Pro Thr Phe Tyr Arg Ser His Tyr Ser Phe Pro Gln Thr Ile Thr His Arg AAA CCC CCA TTT CAT GGC CCT CAC CAT CCT CTT CCC GAC CCA ATA CCA TAC CGC 1251 1260 1269 1278 1287 1296
TTT GGG GGT AAA GTA CCG GGA GTG GTA GGA GAA GGG CTG GGT TAT GGT ATG GCG Phe Gly Gly Lys Val Pro Gly Val Val Gly Glu Gly Leu Gly Tyr Gly Met Ala Leu Gly Val Lys Tyr Arg Glu Trp *** Glu Lys Gly Trp Val Met Val Trp Arg Trp Gly *** Ser Thr Gly Ser Gly Arg Arg Arg Ala Gly Leu Trp Tyr Gly Gly

Pro Thr Thr *** Met Pro Thr Met Pro Ser Pro Gln Pro Arg Gln *** Leu Thr Leu Leu Leu Lys Cys Leu Pro *** Leu His Pro Ser His Gly Lys Asn Cys Leu Ser Ser Tyr Asn Val Tyr Pro Asp Tyr Thr Leu Ala Thr Ala Lys Thr Val Phe CCT CCT CAT CAA ATG TAT CCC CAG TAT CCA CTC CCG ACA CCG GAA ACA ATG TTT 1305 1314 1323 1332 1341 1350 GGA GGA GTA GTT TAC ATA GGG GTC ATA GGT GAG GGC TGT GGC CTT TGT TAC AAA Gly Gly Val Val Tyr Ile Gly Val Ile Gly Glu Gly Cys Gly Leu Cys Tyr Lys Glu Glu *** Phe Thr *** Gly Ser *** Val Arg Ala Val Ala Phe Val Thr Lys Arg Ser Ser Leu His Arg Gly His Arg *** Gly Leu Trp Pro Leu Leu Gln Ser



O.G. FIG.

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ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

Ile Met *** Phe Leu Leu Val Pro Ala Trp Glu Gly Thr Val Arg Pro Ser Arg
*** *** Arg Phe Tyr Cys Cys Gln Leu Gly Ser Gly Gln *** Gly Pro His Asp
Asn Asp Asp Leu Ile Val Ala Ser Ser Gly Val Gly Arg Asp Gly Gln Thr Ile

CAA TAG TAG ATT TTA TTG TCG TGA CCT CGG GTG AGG GGA CAG TGG GAC CCA CTA

1359

GTT ATC ATC TAA AAT AAC AGC ACT GGA GCC CAC TCC CCT GTC ACC CTG GGT GAT

Val Ile Ile *** Asn Asn Ser Thr Gly Ala His Ser Pro Val Thr Leu Gly Asp
Leu Ser Ser Lys Ile Thr Ala Leu Glu Pro Thr Pro Leu Ser Pro Trp Val Ile
Tyr His Leu Lys *** Gln His Trp Ser Pro Leu Pro Cys His Pro Gly *** Ser

Pro Ala Pro Gly Ser Asn Leu Arg Leu Arg Glu *** Glu Thr Thr Asn Leu Pro Pro Leu Leu Ala Leu Ile *** Gly *** Gly Lys Lys Asn Gln Leu Ile *** Leu Pro Ser Cys Pro Trp Phe Glu Val Lys Val Lys Arg Ile Arg Tyr Tyr Glu Phe GCC CCT CGT CCC GGT CTT AAG TTG GAA TTG GAA AGA ATA AGA CAT CAT AAG TTT 1413 1422 1431 1440 1449 1458 CGG GGA GCA GGG CCA GAA TTC AAC CTT AAC CTT TCT TAT TCT GTA GTA TTC AAA Arg Gly Ala Gly Pro Glu Phe Asn Leu Asn Leu Ser Tyr Ser Val Val Phe Lys Gly Glu Gln Gly Gln Asn Ser Thr Leu Thr Phe Leu Ile Leu *** Tyr Ser Lys Gly Ser Arg Ala Arg Ile Gln Pro *** Pro Phe Leu Phe Cys Ser Ile Gln Arg

Cys Leu Ala Pro Thr Gln Gly Gly Glu Gln Pro Phe Phe Thr Met Leu Ile Ser Ala Cys Leu Pro Pro Lys Val Gly Arg Arg Pro Ser Ser Leu *** *** Tyr Gln Pro Val Ser Arg Pro Asn Ser Gly Gly Gly Pro Pro Leu Phe Asp Asn Ile Asn CCC GTG TCT CGC CCC CAA ACT GGG GGG AGG ACC CCC TTC TTT CAG TAA TTA TAA 1467 1476 1485 1494 1503 1512 GGG CAC ACA GCG GGG GTT TGA CCC CCC TCC TGG GGG AGG AAA GTC ATT AAT ATT Gly His Arg Ala Gly Val *** Pro Pro Ser Trp Gly Lys Lys Val Ile Asn Ile Gly Thr Glu Arg Gly Phe Asp Pro Pro Pro Gly Gly Arg Lys Ser Leu Ile Leu Ala Gln Ser Gly Gly Leu Thr Pro Leu Leu Gly Glu Glu Ser His *** Tyr ***

Asp *** *** Thr Trp Arg Gly Pro Pro Arg Glu Ser Gln Pro Glu Ser Ser Leu Ile Glu Asp His Gly Gly Leu Leu Ala Asn Gln Ser His Asn Ala Gln Cys Phe Arg Met Met Asp Val Ala Trp Ser Pro Thr Arg Val Thr Thr Arg Lys Val CCT AGA GTA GTA CAG GTG GCG GGT CCT CCC GCA AGA CTG ACA CCA AGC GAA CTG 1521 1530 1539 1548 1557 1566 GAA TCT CAT CAT GTC CAC CGC CCA GGA GGG CGT TCT GAC TGT GGT TCG CTT GAC Glu Ser His His Val His Arg Pro Gly Gly Arg Ser Asp Cys Gly Ser Leu Asp Asn Leu Ile Met Ser Thr Ala Gln Glu Gly Val Leu Thr Val Val Arg Leu Thr Ile Ser Ser Cys Pro Pro Pro Arg Arg Ala Phe *** Leu Trp Phe Ala *** Gln

Ile Asp Ser Pro Ala Pro Ser Ala Pro Thr Ser Ser Ala Met Lys Gly Glu Gly Tyr Ile Arg Leu His Pro Leu Pro Pro His Gln Leu His Trp Lys Glu Lys Glu Thr Tyr Gly Phe Thr Arg Ser Leu Arg Thr Asn Phe Ile Gly Asn Lys Arg Arg TCA TAT AGG CTT CCA CGC CCT CTC CGC CCA CAA CTT CTA CGG TAA AAA GGA AGA 1575 1584 1593 1602 1611 1620 AGT ATA TCC GAA GGT GCG GGA GAG GCG GGT GTT GAA GAT GCC ATT TTT CTT TCT Ser Ile Ser Glu Gly Ala Gly Glu Ala Gly Val Glu Asp Ala Ile Phe Pro Ser Val Tyr Pro Lys Val Arg Glu Arg Arg Val Leu Lys Met Pro Phe Phe Leu Leu Tyr Ile Arg Arg Cys Gly Arg Gly Gly Cys *** Arg Cys His Phe Ser Phe Ser



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APPROVED | O.G. FIG.



Title: CIRCOVIRUS SEQUENCES
ASSOCIATED WITH PIGLET
WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245



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Ala Thr Val Thr Ala Pro Thr Ser Ser Gly Pro Ala Ala Ala Ser Ser Arg Ala Leu Pro Leu Pro Pro Pro Pro Pro Pro Arg Ala Leu Pro Pro Pro Pro Pro Pro Asp Pro Trp Arg Tyr Arg His Arg Pro His Val Leu Trp Pro Arg Arg Arg Leu Ile Gln GGT CGC CAT TGC CAC CGC CCC CAC CTG CTC GGT CCC CGC CGC CGC CTC CTA GAC 1629 1638 1647 1656 1665 1674 CCA GCG GTA ACG GTG GCG GGG GTG GAC GAG CCA GGG GCG GCG GCG GAG GAT CTG Pro Ala Val Thr Val Ala Gly Val Asp Glu Pro Gly Ala Ala Ala Glu Asp Leu Gln Arg *** Arg Trp Arg Gly Trp Thr Ser Gln Gly Arg Arg Arg Arg Ile Trp Ser Gly Asn Gly Gly Gly Gly Gly Gly Arg Ala Arg Gly Gly Gly Ser Gly

*** Ile Gln Phe Arg Phe Phe His Ala Thr Leu Ile
Asp Tyr Arg Phe Val Phe Ser Thr Arg Gln Leu Tyr
Thr Met Asp Ser Phe Ser Leu Leu Ala Ser Tyr Thr Asn

GCA GTA TAG ACT TTT GCT TTC TTC ACG CGA CAT TCA TAA 5'
1737 1746 1755 1764

CGT CAT ATC TGA AAA CGA AAG AAG TGC GCT GTA AGT ATT 3'

Arg His Ile *** Lys Arg Lys Lys Cys Ala Val Ser Ile
Val Ile Ser Glu Asn Glu Arg Ser Ala Leu *** Val
Ser Tyr Leu Lys Thr Lys Glu Val Arg Cys Lys Tyr

FIG. 8g



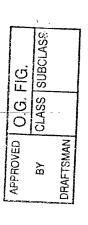
Title: CIRCOVIRUS SEQUENCES
ASSOCIATED WITH PIGLET
WEIGHT LOSS DISEASE (PWD)

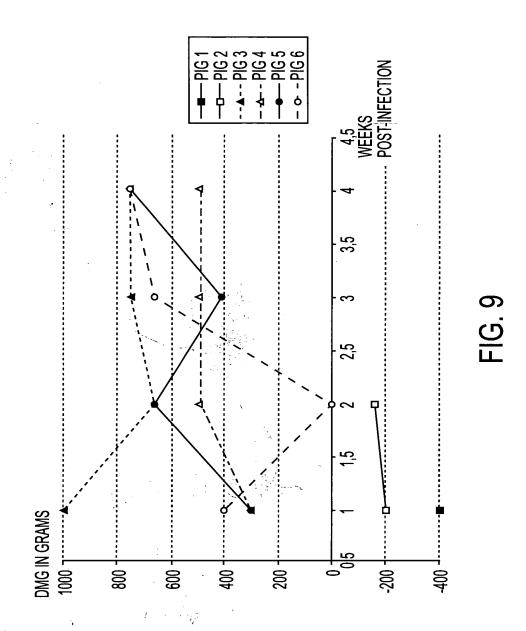
Inventors: Jestin et al. Appl. No.: 09/514,245

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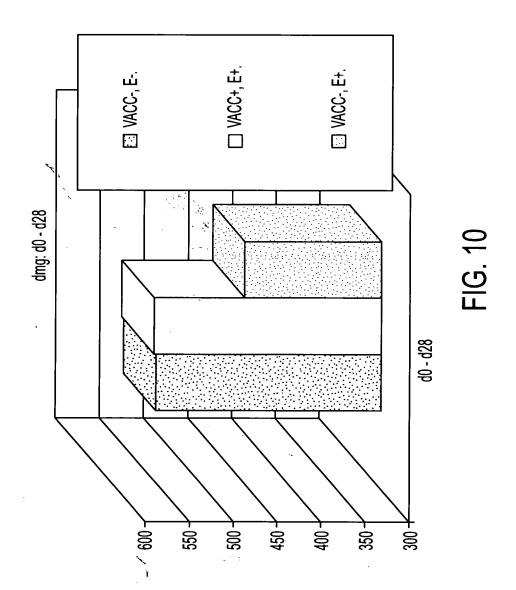
22



Title: CIRCOVIRUS SEQUENCES
ASSOCIATED WITH PIGLET
WEIGHT LOSS DISEASE (PWD)
Inventors: Jestin et al.
Appl. No.: 09/514,245

APPROVED O.G. FIG.

BY CLASS SUBCLASS DRAFTSMAN





Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

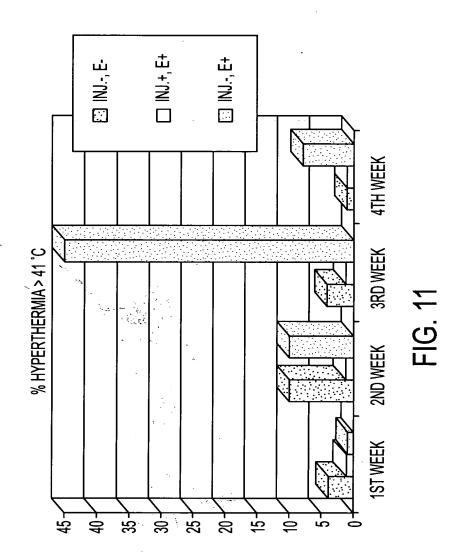
Inventors: Jestin et al. Appl. No.: 09/514,245

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FIG.	SUBCLASS	
O.G. FIG.	CLASS	
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 $(\lambda_{F_{\mathcal{X}_{i}}^{*}}, \omega_{F_{i}}^{*})_{i}$



APPROVED O.G. FIG.

BY CLASS SUBCLASE

DRAFTSMAN

Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245



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FIG. 12

TYPE B SPOT NO. 104 TO 159 TYPE A SPOT NO. 160 TO 215



CLASS SUBCLASS

B

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O.G. FIG.

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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET

WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al. Appl. No.: 09/514,245

152

peptide

RYFTPKPVLD RYFTPKPELD

TIRQPFTYHS TITQPFSYHS

DPYINYSSRH DPYVNYSSRH

pcvA pcvB

151

150

235 P.LNE

ONYVVRLTIY VOFREFILKD

pcvA

pcvB

QEYNIRVTMY VQFREFNFKD

PPLNP

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RGGHSQPSWN RTTVRTPSWA GSTVVILDAN FVTPSTNLAY FVTKATALTY THTNVEHTGL GYALQNATTA TAGNVDHVGL GTAFENSIYD peptide 208 peptides 188 to 189 LSRTFGYTVK GSSAVILDDN LSREFVLTI. to 133 RRKTGIFNSR RRKNGIFNTR QTIDWFQPNN KRNQLWLHLN KRNQLWLRLQ RKAKYEFYPR DPITSNORGV peptides 132 RKVKVEFWPC SPITOGDRGV MTWPRRRYRR RRTRPRSHLG NILRRRPYLV HPAFRNYRW HP. RHYRW FTIDYFOPNN

QILRRRPWLV

RRHRPRSHLG

MTYPRRRYRR

pcvB pcvA

100 LPLPFQYYRI RSVPFEYYRI

VNELRFNIGQ FLPPSGGTNP

pcvA

peptide 177

FLPPGGGSNP

VDMMRFNIND

pcvB

peptide 121



APPROVED O.G. FIG.

BY CLASS SUBCLASS

Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

Inventors: Jestin et al.

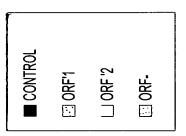
Appl. No.: 09/514,245

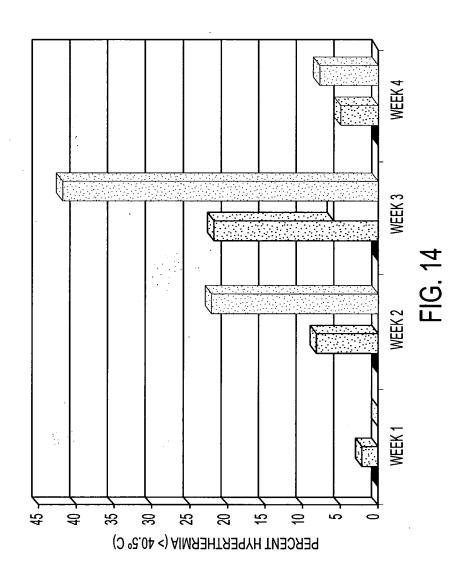


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Title: CIRCOVIRUS SEQUENCES ASSOCIATED WITH PIGLET WEIGHT LOSS DISEASE (PWD)

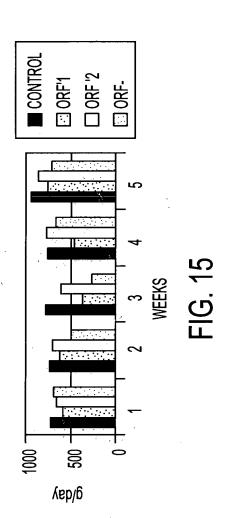
Inventors: Jestin et al. Appl. No.: 09/514,245

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BY CLASS SUBCLASE
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APPROVED O.G. FIG.

BY CLASS SUBCLASS
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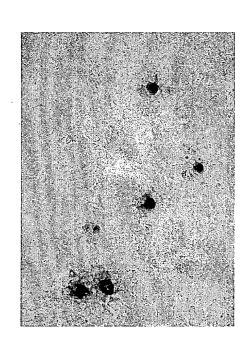


FIG. 16